



**THIS PAGE BLANK (USPTO)**

Attorney Docket No. ORT-1414

**IN THE UNITED STATES PATENT & TRADEMARK OFFICE**

-----X

Re: U.S. PATENT APPLICATION  
Serial Number 09/833,222

Filed: April 11, 2001

Art Unit 1647

-----X

**DECLARATION OF DR. NING QI UNDER RULE 37 CFR 1.131**

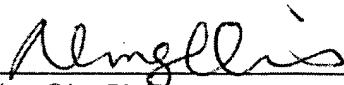
1. I am a co-inventor of the application Serial Number 09/833,222, entitled "cDNA Encoding the Human Alpha2Delta-4 Calcium Channel Subunit" (the '222 application). I obtained my Ph.D. degree in July 1993, from Baylor College of Medicine, and had my postdoctoral training at University of California, Los Angeles from 1994-1997. From 1997 to 1999, I was employed as a research assistant professor at UCLA and then, as an Assistant Professor at University of Kentucky. I have been employed by Johnson & Johnson as a research scientist since September 1999. My resume is attached as Exhibit 1.
2. The invention described in the '222 application is based on the research my colleague and I conducted from 1999 to 2000, during which we successfully cloned the human  $\alpha_2\delta$ -4 calcium channel subunit, as evidenced by the data in my laboratory notebook. I attach the relevant notebook pages (Exhibits 2 and 3 ) and a computer printout (Exhibit 4) herewith to demonstrate the process of our invention. This was completed long before the Brown and Bertelli's publication (March 22, 2001, WO 01/19870A2).

3. I was interested in cloning this subunit because the voltage gated calcium channel is one of important pain therapeutic targets as recognized by the scientists in the field of pain control. I believe that the  $\alpha_2\delta$  is a key regulatory subunit and that there are multiple  $\alpha_2\delta$  isoforms, based on our analysis on human genomic DNA data, all play an important role in mediating  $\text{Ca}^{2+}$  influx in excitable cells.
4. I started the project of cloning  $\alpha_2\delta$ -4 on or about September 20, 1999 (Exhibit 12. No. 15798-1) and my colleague and I continuously worked on the project for the next several months. Because the cDNA encoding  $\alpha_2\delta$ -4 subunit is more than 3Kb, I had to clone it piece by piece. (Exhibit 2). We ultimately assembled and sequenced the fully length cDNA of  $\alpha_2\delta$ -4 subunit by April 7, 2000, which was recorded in my lab notebook (Exhibit 3, p. No. 15798-139), and the DNA and protein sequence information were also documented in both Microsoft word and Seqweb files on the same day. (Exhibit 4).

I declare under the penalty of perjury that the above statements are true and accurate to the best of my knowledge.

Dated: June 6, 2006

By:

  
Ning Qin, Ph.D.

## **CURRICULUM VITAE**

**[04-30-2006]**

**Ning Qin**

**SITE:** Spring House  
**DIVISION:** Analgesics  
**DEPARTMENT:** Drug Discovery  
**EDUCATION:** Baylor College of Medicine, Houston, Texas  
1993, Biochemistry, Ph.D.

### **EMPLOYMENT**

1999-present Johnson & Johnson Pharmaceutical Research and Development  
Spring House, PA  
2003-Present Principal Scientist, Analgesics, Drug Discovery  
1999-2002 Senior Scientist

1999- Department of Pharmacology, University of Kentucky  
Lexington, KY 40536  
1999-1999 Assistant Professor

1994-1998 Dept. of Anesthesiology UCLA. School of Medicine,  
Los Angeles, CA 90095  
1997-1998 Research Assistant Professor  
1994-1996 Postdoctoral Fellow

## CURRENT PROFESSIONAL AFFILIATIONS

1. Society of Neuroscience
2. Biophysical Society
3. American Pain Society
4. International Association for the study of Pain
5. New York Academy of Sciences

## PUBLICATIONS

### *Manuscripts*

1. N. Qin, S. J. Pittler and W. Baehr, (1992) "In Vitro Isoprenylation and Membrane Association of Mouse Photoreceptor cGMP Phosphodiesterase  $\alpha$  and  $\beta$  subunits Expressed in Bacteria", *Journal of Biological Chemistry*, 267, 8458-63
2. N. Qin and W. Baehr, (1993) "Expression of Mouse Rod Photoreceptor PDE  $\gamma$  Subunit in Bacteria", *FEBS Letters*, 321, 6-10
3. M. L. Suber, S. J. Pittler, N. Qin, G. C. Wright, V. Holcombe, R. H. Lee, C. M. Craft, R. N. Lolley, W. Baehr and R. L. Hurwitz, (1993) "Irish Setter Dogs Affected with Rod/Cone Dysplasia Contain a Nonsense Mutation in the Rod cGMP Phosphodiesterase  $\beta$  Subunit Gene", *Proc. Natl. Acad. Sci. U.S.A.* 90, 3968-3972
4. N. Qin and W. Baehr, (1994) "Expression and Mutagenesis of Biologically Active Mouse Rod Photoreceptor cGMP Phosphodiesterase", *Journal of Biological Chemistry*, 269 3265-3271
5. T. Schneider, X. Wei, R. Olcese, J. L. Costantin, A. Neely, P. Palade, E. Peryes, N. Qin, J. Zhou, G. Crawford, R. G. Smith, S. H. Appel, E. Stefani and L. Birnbaumer (1994) "Molecular analysis and functional expression of the human type E neuronal  $\text{Ca}^{2+}$ -channel  $\alpha_1$  subunit", *Receptors and Channels*, 2, 255-270
6. R. Olcese, N. Qin (*Co-first author*), T. Schneider, A. Neely, X. Wei, E. Stefani and L. Birnbaumer (1994) "The Amino Terminus of a Calcium Channel  $\beta$  subunit Sets Rates of Channel Inactivation Independently of the Subunit's Effect on Activation", *Neuron*, 13, 1433-1438
7. F. Noceti, P. Baldelli, X. Wei, N. Qin, L. Toro, L. Birnbaumer and E. Stefani (1996) "Effective Gating Charges Per Channel in Voltage Dependent  $\text{K}^+$  and  $\text{Ca}^{2+}$  Channels", *J. Gen. Physiol.* 108, 143-155
8. N. Qin, R. Olcese, J. Zhou O. Caballo, L. Birnbaumer and E. Stefani (1996) "Identification of a second region of the  $\beta$  subunit involved in regulation of calcium channel inactivation", *Am. J. Physiol. (Cell Physiology)*, 271:C1539-C1545
9. R. Olcese, A. Neely, N. Qin, X. Wei, L. Birnbaumer and E. Stefani (1997) "Coupling Between Charge Movement and Pore Opening in Neuronal  $\alpha_{1E}$  Calcium Channel." *J. Physiol.* 497, 675-686

10. E. Tareilus, M. Roux, **N. Qin**, R. Olcese, J. Zhou E. Stefani and L. Birnbaumer (1997) "A Functional Calcium channel  $\beta$  subunit is Expressed in *Xenopus* Oocytes", *Proc. Natl. Acad. Sci. U.S.A.* 94, 1703-1708
11. J. Zhou R. Olcese, **N. Qin**, E. Stefani and L. Birnbaumer, (1997) "Feedback inhibition of calcium channels by calcium depends on a short sequence of the carboxyterminus that does not include the  $\text{Ca}^{2+}$  binding function of a motif with similarity to calcium binding domains", *Proc. Natl. Acad. Sci. U.S.A.* 94, 2301-2305
12. **N. Qin**, D. Platano, R. Olcese, E. Stefani and L. Birnbaumer (1997) "Inhibition of Neuronal  $\text{Ca}^{2+}$  Channel G-protein Coupled receptors Mediated by a Small C-terminal G $\beta$ y Binding Domain of  $\alpha_1$ ", *Proc. Natl. Acad. Sci. U.S.A.* 94, 8866-887.
13. J. Costantin, F. Noceti, **N. Qin**, X. Wei, L. Birnbaumer and E. Stefani, (1998) "Facilitation by the  $\beta$  Subunit of Pore Openings in Cardiac  $\text{Ca}^{2+}$  Channels", *J. Physiol.* 507, 93-103
14. F. Noceti, R. Olcese, **N. Qin**, and E. Stefani, (1998) " $\text{Ca}^{2+}$ -dependent inactivation in  $\alpha_{1C}$   $\text{Ca}^{2+}$  channels: effect of Bay K 8644(-) and the  $\beta_{2a}$  subunit. A model", *J. Gen. Physiol.* 111, 463-475
15. **N. Qin**, R. Olcese, E. Stefani and L. Birnbaumer (1998) "Modulation of human neuronal  $\alpha_{1E}$  type calcium channel by  $\alpha_2\delta$  subunit", *Am. J. Physiol. (Cell Physiology)*, 274, C1324-C1331
16. **N. Qin**, Platano, R. Olcese, J. Costantin, E. Stefani and L. Birnbaumer (1998) "Unique regulatory properties of type 2a  $\text{Ca}^{2+}$  channel  $\beta$  subunit due to palmitoylation", *Proc. Natl. Acad. Sci. U.S.A.*, 95, 4690-4695
17. N. Li, R.N. Fariss, K. Zhang, A. Otto-Bruc, F. Haeseleer, D. Bronson, **N. Qin**, A. Yamazaki, I. Subbaraya, A. H. Milam, K. Palczewski and W. Baehr, (1998) "Guanylate-cyclase-inhibitory protein is a frog retina  $\text{Ca}^{2+}$  binding protein related to mammalian guanylate-cyclase-activating protein", *Eur. J. Biochem.* 252, 591-59
18. J. Costantin, **N. Qin**, L. Birnbaumer and E. Stefani, (1998) "Long lasting voltage-dependent facilitation of a cardiac and brain calcium channel: correlation with the coupling efficiency between charge movement and pore opening", *FEBS Letter*, 423, 213-217
19. M. Ottolia, D. Platano, **N. Qin** M. Birnbaumer, L. Toro, L. Birnbaumer, E. Stefani and R. Olcese (1998) "Functional Coupling Between Human E-type  $\text{Ca}^{2+}$  Channels and  $\mu$  Opioid Receptors Expressed in *Xenopus* Oocytes"; *FEBS Letter*, 427, 96-102
20. J. Costantin, **N. Qin**, M. N. Waxham, L. Birnbaumer and E. Stefani, (1999) "Complete reversal of run-down in rabbit cardiac  $\text{Ca}^{2+}$  channels by patch-clamping in *Xenopus* oocytes; partial reversal by protein kinase A". *Pflugers Arch*; 437(6): 888-94
21. **N. Qin**, R. Olcese, M. Bransby, T. Lin, and L. Birnbaumer (1999) " $\text{Ca}^{2+}$ -induced inhibition of the cardiac  $\text{Ca}^{2+}$  channel depends on calmodulin". *Proc. Natl. Acad. Sci. U S A*; 96(5): 2435-8
22. B. Vannier, M. Peyton, G. Boulay, D. Brown, **N. Qin**, M. Jiang, X. Zhu, and L. Birnbaumer (1999) "Mouse trp2, the homologue of the human trpc2 pseudogene, encodes mTrp2, a store depletion-activated capacitative  $\text{Ca}^{2+}$  entry channel" *Proc. Natl. Acad. Sci. U S A*; 96(5): 2060-4

23. G. Boulay, D. M. Brown, **N. Qin (Co-first author)**, M. Jiang, A. Dietrich, M. X. Zhu, Z. Chen, M. Birnbaumer, K. Mikoshiba, and L. Birnbaumer (1999) "Modulation of  $\text{Ca}^{2+}$  entry by polypeptides of the inositol 1,4,5-trisphosphate receptor (IP3R) that bind transient receptor potential (TRP): Evidence for roles of TRP and IP3R in store depletion-activated  $\text{Ca}^{2+}$  entry" *Proc. Natl. Acad. Sci. U S A*; 96: 14955-14960.
24. D. Platano, **N. Qin**, F. Noceti, L. Birnbaumer, E. Stefani, and R. Olcese (2000) "Expression of  $\alpha_2\delta$  Subunit Interferes with Prepulse Facilitation in Cardiac L-type Calcium Channels" *Biophys. J.* 78 2959-2972
25. Z. Zhang, J. Tang, S. Tikunova, J. Johnson, Z. Chen, **N. Qin**, A. Dietrich, E. Stefani, L. Birnbaumer, and M. X. Zhu (2001) "Activation of Trp3 by IP<sub>3</sub> Receptors through Displacement of Inhibitory Calmodulin from a Common Binding Domain" *Proc. Natl. Acad. Sci. U S A*; 98: 3168-3173
26. **N. Qin**, S. Yagel, M. Momplaisir, E. E. Codd and M. R. D'Andrea (2002) "Molecular cloning and characterization of human voltage-gated calcium channel  $\alpha_2\delta$ -4 subunit" *Molecular Pharmacology*, 62 (3)
27. J. Leroy, A. Pereverzev, R. Vajna, **N. Qin**, G. Pfitzer, J. Hescheler, C. O. Malécot, T. Schneider, U. Klöckner; (2003) "A novel  $\text{Ca}^{2+}$ -induced regulation of E-type neuronal  $\text{Ca}^{2+}$  channels". *European Journal of Neuroscience*, 18: 841-855
28. **N. Qin**, M. D'Andrea, M. Lubin, N. Shafae, E. Codd and A. Correa (2003) "Cloning and Characterization of a novel splicing variant of human sodium channel  $\beta_1$  subunit". *The FEBS Journal (European Journal of Biochemistry)*, 270: 4762-4770
29. R. Pagani, M. Song, M. McEnery, **N. Qin**, R. W. Tsien, L. Toro, E. Stefani and O. D. Uchitel (2004) "Differential expression of  $\alpha_1$  and  $\beta$  subunits of voltage dependent  $\text{Ca}^{2+}$  channel at the neuromuscular junction of normal and P/Q  $\text{Ca}^{2+}$  channel knockout mouse". *Neuroscience*: 123 (1): 75-85
30. **N. Qin**, SP Zhang, T.L. Reitz, J.M. Mei and C.M. Flores (2005) "Cloning, Expression and Functional Characterization of Human COX-1 Splicing Variants: Evidence for Intron 1 Retention" *The Journal of Pharmacology and Experimental Therapeutics*, 315 (3): 1298-1305
31. Y. Liu, M. Lubin, T.L. Reitz, Y. Wang, R.W. Colburn, C.M. Flores and **N. Qin** (2006) "Molecular identification and functional characterization of a temperature-sensitive transient receptor potential channel (TRPM8) from canine" *European Journal of Pharmacology*, 530: 23-32
32. Y. Liu, **N. Qin**, T.L. Reitz, Y. Wang and C.M. Flores (2006) "Inhibition of the rat brain sodium channel Nav1.2 after prolonged exposure to gabapentin" *Epilepsy Research* 69 (3):

## Book Chapters

1. L. Birnbaumer, **N. Qin**, R. Olcese, E. Tareilus and E. Stefani, (1998). "Studies on the regulation of the human neuronal calcium channel by  $\beta$  and  $\alpha_2\delta$  subunits". *Low voltage activated T-type calcium channels*, 258-268, Montpellier, France

2. F. Noceti, R. Olcese, P. Baldelli, **N. Qin**, L. Birnbaumer and E. Stefani, (1998) "Charges per channel in  $\text{Ca}^{2+}$  channels: effect of the regulatory  $\beta$  subunit co-expression on  $\alpha_1$  pore forming subunit". *Low voltage activated T-type calcium channels*, 307-313, Montpellier, France
3. R. Olcese, F. Noceti, P. Baldelli, **N. Qin**, L. Birnbaumer and E. Stefani, (1998) " $\beta$  subunit modulation of the coupling between charge movement and pore opening in calcium channels". *From Ion Channels to Cell to Cell Conversation*, 91-104, Plenum press, New York and London
4. L. Birnbaumer, **N. Qin**, R. Olcese, E. Tareilus, Daniela Platano and E. Stefani, (1998) Structures and Functions of Calcium Channel  $\beta$  Subunits." *J. of Bioenergetics and Biomembranes*, Vol. 30(4) 357-375
5. L. Birnbaumer, G. Boulay, D. Brown, M. Jiang, A. Doetrich, K Mikoshiba, X. Zhu, and **N. Qin**, (2000) "Mechanism of Capacitative  $\text{Ca}^{2+}$  entry (CCE): Interaction between IP3 Receptor and TRP links the Internal Calcium Storage Compartment to Plasma Membrane CCE Channels." *Recent Progress in Hormone Research*, Vol. 55: 127-162

## ABSTRACTS

1. Y. Liu, M.L. Lubin, T. Reitz, R.W. Colburn, C.M. Flores, and **N. Qin** "Cloning and functional characterization of a canine transient receptor potential channel, cTRPM8". 2004 Neuroscience Meeting
2. **N. Qin**, S-P. Zhang, T. Reitz, E. E. Codd and C. M. Flores "Human COX-3? Cloning and Characterization of a splicing variant of COX-1" 2004 Neuroscience meeting
3. **N. Qin**, M. Neeper, Y. Liu and C. M. Flores "Characterization of calcium channel beta subunit interaction domain" 2005 International Association for the Study of Pain Annual Meeting

## PATENTS

1. **N. Qin**, E. Codd, and M. D'Andrea (2006) "The Human Voltage Gated Sodium Channel  $\beta_{1A}$  Subunit And Method Of Use." *US 6994,993 B2*
2. **N. Qin** and E. Codd (2003) "cDNA encoding the human alpha2delta-4 calcium channel subunit" *US20030166045 A1*
3. **N. Qin**, E. Codd C. M. Flores and SP. Zhang (2004) "Human cyclooxygenase-3 enzyme and uses thereof" *US20040235017 A1*
4. C. M. Flores, Y. Liu, M. Lubin and **N. Qin** (2006) "Canine Cold- And Menthol-Sensitive Receptor.1" *US20060014246 A1*
5. **N. Qin**, Y. Liu, M. Neeper, T. Reitz and C. M. Flores (2005) "Compositions and Methods for Identifying Modulators of TRPV2" Provisional filing
6. **N. Qin**, Y. Liu and C. M. Flores (2005) "A Polypeptide Complex of TRPM8 and Calmodulin And Its Uses Thereof" Provisional filing

RECEIVED 05-2 2001

DOCUMENT ID  
NO. 4368708

BOOK No. 15798



R.W. Johnson Pharmaceutical Research Institute

Issued to Ning Qin

Department Anesthetics Code 62333

Covering the Period 9/7/99 to 6/20/2000

Transferred to \_\_\_\_\_ Date \_\_\_\_\_

R. W. JOHNSON  
PHARMACEUTICAL RESEARCH INSTITUTE  
SPRINGHOUSE

This Document Microfilmed

28 JAN 02 LN-02-01  
(Date) (Film #)

-----  
This notebook is the property of the R.W. Johnson Pharmaceutical Research Institute and constitutes part of its permanent records. It should be returned to the company when not in active use, or upon request.



No. 15798- 1

Project No. X28 of cat channel Protocol / Experiment No. \_\_\_\_\_Date 9/20/99Subject \_\_\_\_\_ Purpose cloning

Contd. from page \_\_\_\_\_

Search Genebank for all published Calcium channels  
916 documents

NCBI Entrez Nucleotide QUERY BLAST Entrez ?

## Current Query

Details

Search : calcium channel[All Fields] --&gt;

Retrieve 916 Documents

Number of documents to display per page: 20 Mod. Date limit: No Limit

## Add Term(s) to Query :

Search Field: All Fields Search Mode: Automatic

Enter Terms: \_\_\_\_\_

Search Clear

Enter one or more author last names, text words, or other keywords. To search for all terms that begin with a given word, place an asterisk (\*) at the end of the word. Journal Titles must be MEDLINE abbreviations; Author names must be in the form LastName Initial(s), e.g. Smith BJ. The initials can be omitted. [Detailed Help](#) is available.

[Clear All](#)

## Modify Current Query :

Term (Total Records)

calcium channel[All Fields] (916)Search for the Intersection (AND) of the selected terms.Questions or comments? Write to the [NCBI Help Desk](#).

Contd. on page \_\_\_\_\_

Investigator Wing QiuDate 9/28/99Read and Understood Ellen CobbDate 100799

No. 15798- 6

Project No.  $\alpha 28$

Protocol / Experiment No. ....

Date 9/28/99

Subject .....

Purpose .....

Contd. from page 5

Analysis and comparison of other small fragments of  $\alpha 28$   
identified on 9/20/99.

Two fragments are novel  $\alpha 28$ .

name them to  $\alpha 28-4$

they are H86016.

J1436938.

Two are human  $\alpha 28-3$  fragments.

they are J3805345

J2885043. or AA001473

sequence comparison results of H86016  
or J1436938

are saved in

H: /mydata/gene / $\alpha 28$  /new- $\alpha 28$ .

Contd. on page 5

Investigator [Signature]

Date 10/1/99

Read and Understood [Signature]

Date 10/7/99

Project No.  $\alpha_2\delta$  Protocol / Experiment No. Date 7/30/99

Subject Purpose

Contd. from page 7

1. comparison of 208672.2  
199408.3

w:66 mouse  $\alpha_2\delta$ -3.

same as H:/mydata/gene/a2d/1nagle clone 208672  
1nagle clone 199408.

2. Design oligos.

A2-3-5

-6

-7

-8

-9

-10

for clone human  $\alpha_2\delta$ -3.

A2-4-1

-2

-3

-4

-5

for clone human  $\alpha_2\delta$ -4.Contd. on page 6 Investigator [Signature] Date 10/1/99Read and Understood E. Cold Date 10/07/99

Project No. d2S E No-31A Protocol / Experiment No. \_\_\_\_\_ Date 10-2-99

Subject \_\_\_\_\_ Purpose \_\_\_\_\_

Contd. from page \_\_\_\_\_

1. Race PCR to amplify NT of h2S-4.  
 Marathon Ready library ① fetal human brain  
 ② human brain.

RX. 5ul 10x Buffer.  
 5ul cDNA library.  
 1ul API.  
 1ul A2-4-9 (10 pmol/ul).  
 1ul Heloanne Tag.  
 1ul 10mM dNTP.  
 35ul H<sub>2</sub>O

cycles. 94°C / 30"  
 ↓  
 94°C / 5" } 5 cycles.  
 ↓  
 72°C / 4 min  
 ↓  
 94°C / 5" } 5 cycles.  
 ↓  
 70°C / 4 min  
 ↓  
 94°C / 5" } 25 cycles.  
 ↓  
 68°C / 4 min  
 ↓  
 4°C / parking.

Contd. on page 22Investigator [Signature]Date 11/16/99Read and Understood [Signature]Date 3-6-00

Project No. 225 + Na-β1A Protocol / Experiment No. \_\_\_\_\_ Date 11-89

Subject \_\_\_\_\_ Purpose \_\_\_\_\_

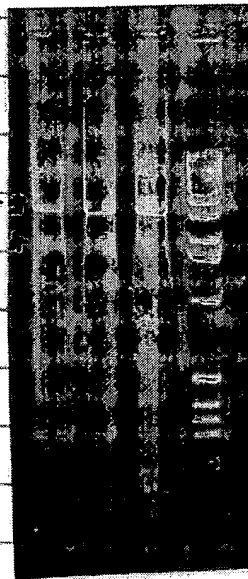
Contd. from page \_\_\_\_\_

1. cloning Race PCR products (11-5-99) of NT α2S-4 (human)

① gel purification  
cut two parts.

A. 3 Kb band

B 2-2.5 Kb bands

purify them  
separately

②. polish them.

③ Ligation: pScript.

④ transfer to XL10-gold

2. Retest condition for Na-β1A.

RX 50 μl, with Advantage Tag

① SB1-4 + SB1-2

② SB1-5 + SB1-2

③ SB1-1 + SB1-2

④ SB1-5 + SB1-2

} rat brain cDNA

} human brain cDNA

condition: 30 cycles.

94°C/30" → 58°C/30" → 72°C/60"

5 μl to check PCR RX.

does not work.

Contd. on page 25Investigator [Signature]Date 11/16/99Read and Understood [Signature]Date 3-6-00

Project No. d<sub>2</sub>8. Protocol / Experiment No. \_\_\_\_\_ Date 11-11-99

Subject \_\_\_\_\_ Purpose \_\_\_\_\_

Contd. from page \_\_\_\_\_

1. miniprep NQC44.  
 two new 3kb  $\alpha$ S-4 in PCR2.1  
 four new 2kb  $\alpha$ S-4 in PCR2.1

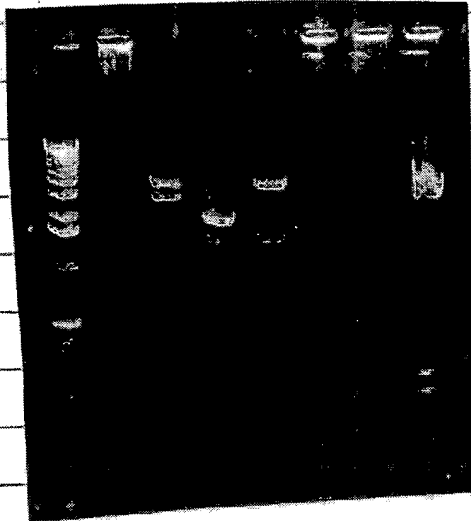
dig. NQC44 with EcoRI + KpnI  
 all others with EcoRI only

3-1 3-2 2-1 2-2 3-2 4, NQC44

3-2, 2-1, & 2-2  
 have insertion

3-2  $\rightarrow$  NQC45  
 2-1  $\rightarrow$  NQC46  
 2-2  $\rightarrow$  NQC47

Send NQC44-47  
 for ~~seq~~

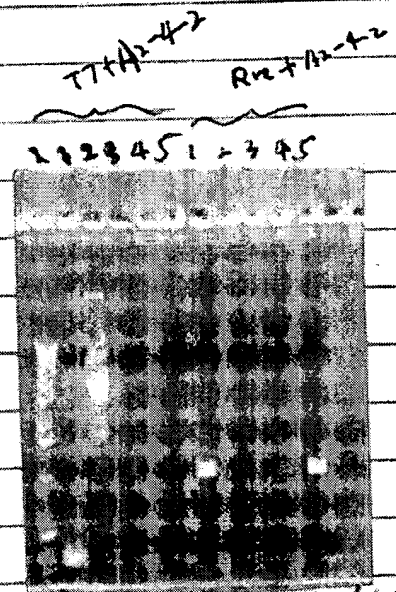


2. characterise these clones. by PCR.  
 ① NQC44, ② NQC45, ③ NQC46  
 ④ NQC47, ⑤ lysate 3343341

20  $\mu$ l RX.

NQC44 & 45 are very promising  
 but 44 is too short.

45 is right size.

Contd. on page 32Investigator ASRead and Understood ni/wDate 11/16/99Date 3-6-00

Project No. α<sub>2</sub>S Protocol / Experiment No. \_\_\_\_\_ Date 11-11-99

Subject \_\_\_\_\_ Purpose \_\_\_\_\_

Contd. from page \_\_\_\_\_

1. miniprep NQC44.  
two new 3kb α<sub>2</sub>S-4 in PCR2.1  
four new 2kb α<sub>2</sub>S-4 in PCR2.1

dig. NQC44 with EcoRI + KpnI  
all others with EcoRI only

3-1 3-2 2-1 2-2 2-3 2-4, NQC44.

3-2, 2-1, & 2-2  
have insertion

3-2 → NQC45  
2-1 → NQC46  
2-2 → NQC47

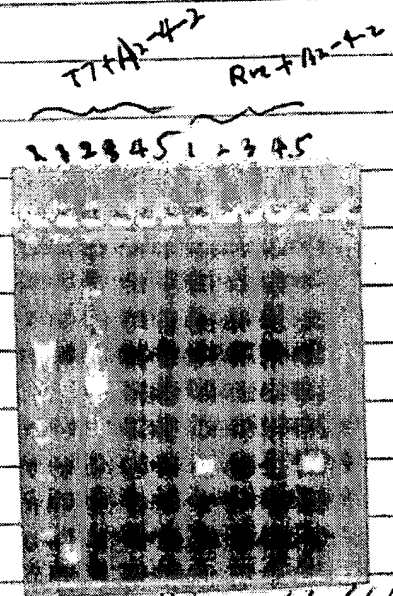
Send NQC44-47  
for seq.



2. characterise these clones. by PCR.  
① NQC44, ② NQC45, ③ NQC46  
④ NQC47, ⑤ lysate 3343341

20 ul RX.

NQC44 & 45 are very promising  
lab 44 is too short.  
45 is right size.



Contd. on page 32

Investigator \_\_\_\_\_

Read and Understood \_\_\_\_\_

Date 11/16/99

Date 3-6-00

Project No. Wn-β / Cu-ds8 Protocol / Experiment No. \_\_\_\_\_ Date 4/7/00  
Subject \_\_\_\_\_ Purpose \_\_\_\_\_

Contd. from page \_\_\_\_\_

2. Analysis of ~~ss~~ data of hds-4 NT.  
Thay (NAC 190-195) are all hds-4 NT.  
finished cloning of ds-4.

Syn oligos: A2-4-27 } for add HA on hds-4 CT  
A2-4-28 }

A2-3-29 } add HA on  
A2-3-30 } hds-3 CT.

Contd. on page 137

Investigator [Signature]

Date 4/14/00

Read and Understood [Signature]

Date 4/17/00



**ha2d-4b.pep**[Text View](#)

!!AA\_SEQUENCE 1.0

ha2d-4.pep Length: 1090 April 7, 2000 14:19 Type: P Check: 9269 ..

```
1  MAVALGTRRR DRVKLWADTF GGDLYNTVTK YSGSLLLQKK YKDVESSLKI
51  EEVDGLELVR KFSEDMENML RRKVEAVQNL VEAAEEADLN HEFNESLVFD
101 YNSVLINER DEKGNFVELG AEFLLESNAH FSNLPVNTSI SSVQLPTNVY
151 NKDPDILNGV YMSEALNAVF VENFQRDPTL TWQYFGSATG FFRIYPGIKW
201 TPDENGVITF DCRNRGWYIQ AATSPKDIVI LVDVSGSMKG LRMTIAKHTI
251 TTILDTLGEN DFNIIAYND YVHYIEPCFK GILVQADRDN REHFKLLVEE
301 LMVKGVGVD QALREAFQIL KQFQEAQGS LCNQAIMLIS DGAVEDYEPV
351 FEKYNWPDCK VRVFTYLIGR EVSFADRMKW IACNNKGYTT QISTLADTQE
401 NVMEYLHVLS RPMVINHDHD IIWTEAYMDS KLLSSQAQSL TLLTTVAMPV
451 FSKKNETRSH GILLGVVGS VALRELMKLA PRYKLGVHGY AFLNTNNGYI
501 LSHPDLRPLY REGKKLKPKP NYSVDLSEV EWEDQAESLR TAMINRETGT
551 LSMDVKVPMD KGKRVFLTN DYFFTDISDT PFSLGAVLSR GHGEYILLGN
601 TSVEEGLHDL LHPDLALAGD WIYCITDIDP DHRKLSQLEA MIRFLTRKDP
651 DLECDEELVR EVLFDVVTA PMEAYWTALA LNMSESEHV VDMAFLGTRA
701 GLLRSSLFVG SEKVSDRKFL TPEDEASVFT LDRFPLWYRQ ASEHPAGSFV
751 FNLRWAEQPE SAGEPMVVTA STAVAVTVDK RTAIAAAAGV QMKLEFLQRK
801 FWAATRQCST VDGPYTQSCD DSDLDCFVID NNGFILISKR SRETGRFLGE
851 VDGAVLTQLL SMGVFSQVTM YDYQAMCKPS SHHHSAAQPL VSPISAFLLA
901 TRWLLQELVL FLLEWSVWGS WYDRGAEAKS VFHSHKHKK QDPLQPCDTE
951 YPVFVYQPAI REANGIVECG PCQKVFFVQQ IPNSNLLLLV TDPTCDCSIF
1001 PPVLQEATEV KYNASVKCDR MRSQKLRRRP DSCHAFHPEV RVEADRGWAG
1051 FSSPNPLCLG LCPCRQEHIG MPMNTPVPVL LGGNIRVYAL
```

## Human $\alpha_2\delta_4$ map

# HindIII

AGTCTGCCACTCTCCAAACCcAGAGGCCCTTGGGCTCAAGCTCAGTCTTGGGCTCGTCAGCCC  
1 - - - - - + - - - - - + - - - - - + - - - - - + - - - - - + - - - - - + - - - - - +  
TCAGACGGTGAGAGGTTGGTCTCCGGGACCTTCGAACCCcAGTTCAGTCAGGACCCGAGCAGTCCGGG  
90

B5751

[illegible]

NcoI Splicing

A2-4-37, -39 (*Kozak & HindIII*)  
 GCCCAGGCCATGGCTGTAGCTTTAGGGACAAGGAGGAGGGACAGAGTGAAGCTATGGGCTGACACCTTCGGCGGGACCTGTATATAACACT  
 181  
 CGGGTCCGGTACCGACATCGAAATCCCTGTTCCTCCTCCCTGTCTCACTTCGATACCCGACTGTGGAAAGCCGCCCTGGACATATTGTGA  
 M A V A L G T R R R R D R V K L W A D T F G G D L Y N T -  
 270

## PsTI

BspGI

271 GTGACCAAATACTCAGGCTCTCTCTGTGCAGAAAGTATCAAGGATGTGGAGTCCAGTCTGAAGATCGAGGAGGTGGATGGCTTGGAG  
 -----+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+  
 CACTGGTTATGAGTCCGAGAGAGAAACGACGTCTTCTTCATGTCTCTACACCTCAGGTCAGACTTCTAGCTCCTCCACCTACCGAACCTC  
 V T K Y S G S L L L L O K K Y K D V E S S L K I E E V D G L E -  
 360

BspGI

[illegible]

## ECORI

DrdII

GACCTGAACCAAGAAATTCCTGGTTCGACTATTACAACTCGGTCTGATCAACGAGGGGACGACGAGGGGACGCTTCG  
 451  
 CTGGACTTGGTCTTAAGTTACTTAGGGACCACAAGCTGAATATGTTGAGCCAGGACTAGTTGCTCTCCCGTCTTCCCGTGAAGCAC  
 540  
 D L N H E F N E S L I V F D Y Y N S V L I N E R D E K G N F V -

NarI | PvuII  
 A2-4-20  
 541 GAGCTGGGCGCGAGTTCCTCTGGAGTCAATGCTCaCTTCAGCAACCTGCGCGTGAACACCTcCATCAGCAGCGTGCAGCTGCCACC  
 CTGACCCCGCGCTCAAGGAGGACCTCAGGTTACGAGTCAAGTCTGGACGGCCaCTTGTGGAGGTAGTCGTGCACGTGCACGGGTGG  
 E L G A E F L L E S N A H F S N L P V N T S I S S V Q L P T - 630  
 A2-4-24  
 BsrGI | DraI | PciI | BsmI  
 631 AACGTGTACAAAGAGCCAGATATTTTAAATGGAGTCTACATGTCTGAAGCCTTGAATGCTGTCTTCGTGGAGAACTTCAGAGAGAC  
 TTGCACATGTTGTTCTGGGTCTATAAAATTTACCTCAGATGTACAGACTTCGGAACCTTACGACAGAAACCTCTTTGAAGGTCTCTCTG  
 N V Y N K D P D I L N G V Y M S E A L N A V F V E N F Q R D -  
 A2-4-25  
 SspI | BtsI  
 721 CCAACGTTGACCTGGCAATATTTTGGCAGTGCACACTGGATTCTTCAGGATCTATCCAGGTATAAAATGGACACCTGaTGAGAAATGGAGTC  
 GGTTCACACTGGACCGTTATAAAACCGTCACGTTGACCTAAGAGTCTCTaGATAGGTCCATATTTTACCTGTGGACTACTCTTACCTCAG  
 P T L T W Q Y F G S A T G F F R I Y P G I K W T P D E N G V - 810  
 A2-4-19  
 SacII | BsrBI  
 811 ATTACTTTGACTGCCGAAACCGCGGCTGGTACATTCAAGCTGCTACTTCTCCAAAGACATAGTGAATTTTGGTGGACGTGAGCGGCAGT  
 TAATGAAAACCTGACGGCTTTGGCGCGACCATGTAAGTTTCGACGATGAAGAGGGTTCCTGTATCACTAAACACCTGCACCTGCCGTCA  
 I T F D C R N R G W Y I Q A A T S P K D I V I L V D V S G S - 900  
 A2-4-26  
 BamHI | DraI | BspGI  
 901 ATGAAGGGCTGAGGATGACTATTGCCaAGCACAACCACCATCTTGGACACCCCTGGGGAGAAATGACTTCGTTAATATCATAGCG  
 TACTTCCCGACTCCTACTGATAACGGTTCGTGTGGTAGTGGTAGAACCTGTGGGACCCCTCTTACTGaAGCAATTATAGTATCGC  
 M K G L R M T I A K H T I T T I L D T L G E N D F V N I I A -  
 A2-4-32  
 991 TACAATGACTACGTCCATTACATCGAGCCCTTTTAAAGGATCCTCGTCCaGGCGGACCGAGACAATCGACAGCATTTCAAACCTGCTG  
 ATGTTACTGATGACGTAATGATGCTCGGAACAAAATTTCCCTAGGACGAGTCCGCGCTGGCTCTGTAGCTCTCGTAAAGTTTGACGAC  
 Y N D Y V H Y I E P C F K G I L V Q A D R D N R E H F K L L -  
 A2-4-32  
 1081 GTGGAGGAGTTGATGTCAAAAGTGTGGGGTCTGTGGACCAAGCCCTGAGAGAAGCCTTCCAGATCTCGAAGCAGTTCCAGAGGCCAAG  
 CACCTCCTCAACTACAGTTTCACACCCCCAGCACCTGGTTCCGGAGGTCTCTTCCGGAGGTCTAGGACTTCGTAAGGTCTCCTCGGTTC  
 V E L M V K G V G V D Q A L R E A F Q I L K Q F Q E A K - 1170  
 NarI | Pfl11081 | MscI

1171 CAAGGAAGCTCTGTGCAACGAGGCCATCATGCTCATCAGCGACGCGCGGTGAGGACTACGAGCCGGTGTGAGAAAGTATAACTGGCCA  
 1260  
 a GTTCCTTCGGAGACGTTGGTCCGGTAGTACGAGTAGTCGTCGCGGCACCTCCTGATGCTCGGCCACAACTCTTCATATTGACGGGT  
 Q G S L C N O A I M L I S D G A V E D Y E P V F E K Y N W P  
 A2-4-13 A2-4-12 A2-4-11  
 SphI

1261 GACTGTAAAGTCCGAGTTTCACTTACTTCACTTCACTTGGGAGAGAAAGTGTCTTTTGTGTGACCGCATGAAGTGGATTGCATGCAACAAAGGC  
 1350  
 a CTGACATTCAGGGCTCAAAAGTGAATGGAGTAACCCCTCTCTTCACAGAAAAACGACTGGCGTACTTCACTAACGTACGTTGTTGTTCCG  
 D C K V R V F T Y L I G R E V S F A D R M K W I A C N N K G

1351 tACTACAGCAGATCTCAACGCTGGCGGACACCCAGGAGAACGTGATGAATACCTGCACGTGCTCAGCCGCCCATGGTTCATCAACCCAG  
 1440  
 a ATGATGTGCGTCTAGAGTTGCGACCGCTGTGGTCTCTTGGCACTACCTTATGGACGTGCACGAGTCGGCGGGGTACCCAGTAGTTGGTG  
 Y Y T Q I S T L A D T Q E N V M E Y L H V L S R P M V I N H  
 BspGI StuI BtsI MscI  
 A2-4-29 A2-4-18 A2-4-17  
 A2-4-16

1441 GACCACGACATCATCTGGACAGAGGCCCTACATGGACAGAAAGCTCCTCAGCTCGAGGCTCAGAGCCTGACACTGTCACTGTGGCC  
 1530  
 a CTGGTGTGTAGTACCTGTCTCCGGATGTACCTGTCTCGAGAGTCGAGCTCCGAGTCTCGAGTGTGACGAGTGGTGACACCGG  
 D H D I I W T E A Y M D S K L S S O A O S L T L L T T V A  
 NcoI BsmI

1531 ATGCCAGTCTTCAGCAAGAAGAACGAAACGCGATCCCATGGCAATCTCCTGGTGTGGTGGCTCAGATGTGGCCCTGAGAGAGCTGATG  
 1620  
 a TACGGTCAGAAGTCGTTCTTGTGCTTTGCGTAGGGTACCGTAAGAGGACCCACACCCGAGTCTACACCGGGACTCTCTCGACTAC  
 M P V F S K K N E T R S H G I L L G V V G S D V A L R E L M  
 NarI HindIII  
 AAGCTGGCGCCCGGTACAAGCTTGGAGTGCACGGATACGCCTTTCTGAACCAACAATGGCTACATCTCTCCCATCCGACCTCCGG  
 1621  
 TTCGACCGCGGGCCCATGTTTCGAACCTCAAGTCCGCTATGCGGAAAGACTTTGTGGTTGTACCGATGTAGGAGAGGGTAGGGCTGGAGGCC  
 K L A P R Y K G V H G Y A F L N T N N G Y I L S H P D L R  
 A2-4-30

BsrGI

1711 CCCCTGTACAGAGGGGAGAACTAAACCCAAACCTAACTAACAGTGTGGATCTCTCCGAAGTGGAGTGGGAAGACCAGGCTGAA 1800

a P L Y R E G K K L K P K P N Y N S V D L S E V E W E D Q A E -

1801 TCTCTGAGAACAGCCATGATCAATAGGGAACACAGGTACTCTCTCGATGGATGTGAAGTTCCGATGGATAAAGGAAGCGAGTCTTTTTC 1890

a S L R T A M I N R E T G T L S M D V K V P M D K G K R V L F -

SmaI

1891 CTGACCAATGACTACTTCTTACGGACATCAGCGACACCCCTTTTCAGTTTGGGGTGGTGTCTGTCCCGGGGCCACGGAGAATACATCCCTT 1980

a L T N D Y F F T D I S D T P F S L G V V L S R G H G E Y I L -

BtrI StuI

1981 CTGGGGAACACGTCGTGGAAGAAGCCCTGCATGACTTGTCTTACCCAGACCTGGCCCTGGCCGGTGAATGATCTACTGCATCACAGAT 2070

a L G N T S V E E G L H D L L H P D L A L A G D W I Y C I T D -

2071 ATTGACCCAGACCCGGAAGCTCAGCCAGCTAGAGGCCATGATCCGCTTCTCACCAGGAAGACCAGACCTGGAGTGTGACGAGGAG 2160

a I D P D H R K L S Q L E A M I R F L T R K D P D L E C D E E -

NcoI

BspGI

PciI

2161 CTGCTCCGGGAGGTGCTGTTTGACGCGGTGGTGACAGCCCCCATGGAAAGCCTACTGGACAGCGCTGGCCCTCAACATGTCCGAGGAGTCT 2250

a L V R E V L F D A V V T A P M E A Y W T A L A L N M S E S -

PmlI

SmaI

2251 GAACACGTGGTGACATGGCCCTTCTGGGCACCCCGGGCTGGCCCTCCTGTAGAAGCAGCTTGTTCGTGGGCTCCGAGAAAGTCTCCGACAGG 2340

a CTTGTGCACCACTGTACCGGAAGGACCCGTGGGCCCGACCGGAGGACTCTTCGTGCAACAAGCACCCGAGGCTCTTCCAGAGGCTGTCC K V S D R -

**A2-4-34**

a AAGTTCCTGACACCTGAGGACGAGGCCAGCGTGTTCAACCTGGACCGCTTCCCGCTGTGGT GCCAGGCCTCAGAGCATCTCTGCTGSC  
+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+  
2341 TTCAAGGACTGTGACTCTCTGCTCCGGTCGACAAGTGGACCTGGCGAAGGCGCACACCATGGCGGTCCGGAGTCTCGTAGGACGACCG  
K F L T P E D E A S V F T L D R F P L W Y R Q A S E H P A G -  
**Antigen1**

NcoI PvulI StuI

a AGCTTCGTCTTCAACCTCCGCTGGCAGAACGACCAAGTGGGGTGAACCCATGGTGTGTGACGGCAAGCACAGCTGTGGCGGTGACC  
+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+  
2431 TCGAAGCAGAAGTTGGAGGCGACCCGCTCTTCTGGTCTTTTCAAGCCCACTTGGGTACCACTGCGGTTTCGTGTGACACCGCCCACTGG  
S F V F N L R W A E G P E S A G E P M V V T A S T A V A V T -  
BspGI EcoRI BtsI

a GTGACAAGAGGACAGCCATTGCTGCAGCCGCGGGCGTCCAATGAAGCTGGAATTCCTCCAGCGCAAATTCCTGGCGGCAACGCGGCAG  
+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+  
2521 CACCTGTTCTCCTGTTCGGTAACGACGTCGGCGCCCGCAGGTTTACTTCGACCTTAAGGAGTGGCGTTTAAGACCCGCGGTTGCGCCGTC  
V D K R T A I A A A A G V Q M K L E F L R K F W A A T R Q -  
**A2-4-2**

BspGI

a TGCAGCACTGTGGATGGGCGGTGCACACAGAGCTGCGAGGACAGTGATCTGGACTCTTCGTCATCGACAACAACGGGTTTCATTCTGATC  
+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+  
2611 ACGTCGTGACACCTACCCGGCACGTGTCTCGACGCTCCTGTCTACCTAGACCTGACGAAGCAGTAGCTGTTGCCCCAAAGTAAGACTAG  
C S T V D G P C T Q S C E D S D L D C F V I D N N G F I L I -  
**A2-4-7**

PvuII

a TCCAAGAGGTCCCAGAGACGGAAGATTCTTGGGGAGGTggaTGGTGTGTCTCTGACCCAGCTGCTCAGCATGGGGGTGTTTCAGCCAA  
+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+  
2701 AGGTTCTCCAGGGTCTCTGCCCCCTTCTAAAGACCCCCCTCCACctACCACGACAGGACTGGGTTCGACGAGTCGTACCCCCCAAGTCGGTT  
S K R S R E T G R F L G E V D G A V L T Q L L S M G V F S Q -  
**A2-4-10, -3 (sh)**

**A2-4-42**

a GTGACTTATGACTATCAGGCCATGTGCAAA TCGAGTCACCACCACAGTCGAGCCCCAGCCCCCTGGTCAGCCCCAATTTCTGCCTTC  
+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+  
2791 CACTGATACATACTAGTACCGGTACACGTTTGGGAGCTCAGTGGTGTGTACGTTCGGGTTCGGGACCCAGTCGGGTTAAAGACGGAAG  
V T M Y D Y Q A M C K P S S H H S A A O P L V S P I S A F -  
**A2-4-9, -4 (Sh)**

PstI

2881 TTGACGGCGACAGGTGGCTGCTGCAGGAGTGTGTGTTCTGCTGCTGAGTGTGAGTGTCTGGGGCTCTGGTACGACAGAGGGGCGcgaG

a AACTGCCGCTGTCTCCACCGACGACGTCTCTGACCAACAGACCACTCACTCACAGACCCCGAGGACCACTGTGTCTCCCCGGgtc  
L T A T R W L L Q E L V L F L L E W S V W G S W Y D R G A E -

PstI

NgoAIV

2971 GCCAAAAGTGTCTTCCATCACTCCACAAACAAaagaagCAGGACCCGCTGcagcCCTGcagacaCGgagtACCCcgTGTtcGTGTAccaG

a CGGTTTTTCACAGAAGGTAGTGGGTGTTGTGTTcttcGTCTCGGGCGACgtcgGGACgctgtGCctcaTGGggcACAaagCACATggtc  
A K S V F H S H K H K K Q D P L Q P C D T E Y P V F V Y Q -

ApaI

3061 CCGGccaTCCGGGagGCCAACGGGATCGTGGAGTGGGGCCCTGTGCAGAAAGTATTTGTGTGTCAGCAGATTCCTCAACAGTAACCTCCTC

a GGCCggtAGGCCctccGGTTGCCCTAGCACCTCACGCCCGGGACGGTCTTCCATAAAACACCACGTCTGTCTAAGGGTTGTTCATTGGAGGAG  
P A I R E A N G I V E C G P C Q K V F V V Q Q I P N S N L L -

PstI

PstI

A2-4-33

3151 CTCCTGTGTACAGACCCACCTGTGACTGCAGCATCTTCCACCAGTGTGCAGGAGGGCGACAGAAAGTCAAAATATAATGCTCTGTCAAA

a GAGGACCACTGTCTGGGTGGACACTGACGTCTGTAGAGGGTGTACGACGTCTCCGCTGTCTTCAAGTTTATATTACGGAGACAGTTT  
L L V T D P T C D C S I F P P V L Q E A T E V K Y N A S V K -

NgoAIV

Splicing

3241 TGTGACCGGATCGCTCCAGaagctccGCCGGGACACAGACTCCTGCCACGCCTTCCATCCAGAGTGTGGGGTTGAGCGGATCGAGGG

a ACCTGGCCTACGCGAGGTcttcgagCGGCCGCTGCTGAGGACGGTGGGAAGGTAGGTCTCCACGCCCAACTCCGCTAGCTCCC  
C D R M R S Q K L R R R P D S C H A F H P E V R V E A D R G -

Antigen2

PstI

3331 TGGGCTGGATTTTCATCCCCAAACCCTCTGTGCTGTGGTCTGTGCCCTGCAGACAGGAGCATATAGGGATGCCAATGAACACACCTGTG

a ACCCGACCTAAAGTAGGGTGTGGGAGACACCGGACCCAGACACGGGGACGTCTGCTCCTGATATATCCCTACGGTTACTTGTGTGGACAC  
W A G F S S P N P L Q L G L C P C R Q E H I G M P M N T P V -

CCTGTGCTTCTCGGGGAAACATTCCGGTTTTATGCCCTGTGACACTGTGATATAATAAGAAACAGA

3421 GGACACGAAGAGCCCCCTTTGTAAGCGCAATACGGGACACTGTGACACTATATATCTTTCTCT

a P V L L G G N I R V Y A L \* A2-4-38, -8 A2-4-28 (HA) □

**Enzymes that do cut and were not excluded:**

Apal	BamHI	BglII	BsmI	BspGI	BsrBI	BsrGI	BtrI	BtsI	DraI	DrdII	EcoRI
HindIII	KpnI	MscI	NarI	NcoI	NgoAIV	PciI	Pfl1108I	PmlI	PstI	PvuII	SacII
SmaI	SphI	SspI	StuI	XhoI							

**Enzymes that do not cut:**

Bsbl	BssHII	BssSI	BstZ17I	Clal	EagI	EcoRV	HpaI	Kpn2I	MluI	MunI	NdeI
NheI	NotI	NruI	NsiI	NspV	PacI	PirAI	PsiI	PvuI	RcaI	SacI	Sall
ScaI	SnaBI	SpeI	SrfI	SunI	VspI	XbaI					



**DNA Sequence:**

```
1 CAGGTACATT CAGCAGAGCC CAAGTCTGCC ACTCTCCAAC CaGAGGCCCT
51 GGAAGCTTGG GGTCAAGCTC AGTCCTGGGC TCGTCAGCCC GGCCCCACAA
101 CCCTCAGCAG GAGaACCTGC CGAGGACATT CAGCACACAG CAGTGCAGCC
151 GCTGGGTCCT GAGGGTTCTC CGCGTCTCCT GCCCAGGCCA TGGCTGTAGC
201 TTTAGGGACA AGGAGGAGGG ACAGAGTGAA GCTATGGGCT GACACCTTCG
251 GCGGGGACCT GTATAACaCT GTGACCAAAT ACTCAGGCTC TCTCTTGCTG
301 CAgAAGAAGT ACAAGGATGT GGAGTCCAGT CTGAAGATCG AGGAGGTGGA
351 TGGCTTGGAG CTGGTGAGGA AGTTCTCAGA GGACATGGAG AACATGCTGC
401 GGAGGAAAGT CgAGGCGGTC CAgAATCTGG TGGAAAGCTGC CGAGGAGGCC
451 GACCTGAACC ACGAATTCAA TGAATCCCTG GTGTTCTGACT ATTACAACCTC
501 GGTCCTGATC AACGaGAGGG ACGAGAAGGG CaACTTcGTG GAGCTGGGCG
551 CCGAGTTCCT CCTGGAGTCC AATGCTCaCT TCAGCAACCT GCCGGtGAAC
601 ACCTcCATCA GCAGCGTGCA GCTGCCACC AACGTGTACA ACAAAGACCC
651 AGATATTTTA AATGGAGTCT ACATGTCTGA AgCCTTGAAT GCTGTCTTCG
701 TGGAGAACTT CCAGAGAGAC CCAACGTTGA CCTGGCAATA TTTTGGCAGT
751 GCAACTGGAT TCTTCAGGAt CTATCCAGGT ATAAAATGGA CACCTGaTGA
801 GAATGGAGTC ATTACTTTTG ACTGCCGAAA CCGCGGCTGG TACATTCAAG
851 CTGCTACTTC TCCCAAGGAC ATAGTGATTT TGGTGGACGT GAGCGGCAGT
901 ATGAAGGGGC TGAGGATGAC TATTGCCaAG CACaCCATCA CCACCATCTT
951 GGACACCCTG GGGGAGAATG ActTCGTTAA TATCATAGCG TACAATGACT
1001 ACGTCCATTA CATCGAGCCT TGTTTTAAAG GGATCCTCGT CCaGGCGGAC
1051 CGAGACAATC GAGAGCATTT CAAACTgCTG GTGGAGGAGT TGATGGTCAA
1101 AgGTGTGGGG GTCGTGGACC AAGCCCTGAG AGAAGCCTTC CAGATCCTGA
1151 AgCAGTTCCA AGAgGCCAAG CAAGGAAGCC TCTGCAACCA GGCCATCATG
1201 CTCATCAgCG ACgGCGCCGT GGAGGACTAC GAGCCGGTGT TTGAGAAGTA
1251 TAACTGGCCA GACTGTAAGG TCCGAGTTT CACTTACCTC ATTGGGAGAG
1301 AAGTGTCTTT TGCTGACCGC ATGAAGTGGA TTGCATGCAA CAACAAAGGc
1351 tACTACACGC AGATCTCAAC GCTGGCGGAC ACCCAGGAGA ACGTGATGGA
1401 ATACCTGCAC GTGCTCAGCC GCCCATGGT CATCAACCAC GACCACGACA
1451 TCATCTGGAC AGAGGCCTAC ATGGACAGCA AGCTCCTCAG CTCGCAGGCT
1501 CAGAGCCTGA CACTGCTCAC CACTGTGGCC ATGCCAGTCT TCAGCAAGAA
1551 GAACGAAACG CGATCCCATG GCATTCTCCT GGGTGTGGTG GGCTCAGATG
1601 TGGCCCTGAG AGAGCTGATG AAGCTGGCGC CCCGGTACAA GCTTGGAGTG
1651 CACGGATACG CCTTTCTGAA CACCAACAAT GGCTACATCC TCTCCCATCC
1701 CGACCTCCGG CCCTGTACA GAGAGGGGAA GAAACTAAAA CCCAAACCTA
1751 ACTACAACAG TGTGGATCTC TCCGAAGTGG AGTGGGAAGA CCAGGCTGAA
1801 TCTCTGAGAA CAGCCATGAT CAATAGGGAA ACAGGTAActC TCTCGATGGA
1851 TGTGAAGGTT CCGATGGATA AAGGGAAGCG AGTTCTTTTC CTGACCAATG
1901 ACTACTTCTT CACGGACATC AGCGACACCC CTTTCAGTTT GGGGGCGGTG
1951 CTGTCCCGGG GCCACGAGA ATACATCCTT CTGGGGAACA CGTCTGTGGA
2001 AGAAGGCCTG CATGACTTGC TTCACCCAGA CCTGGCCCTG GCGGTGACT
2051 GGATCTACTG CATCAGATG ATTGACCCAG ACCACCGAA GCTCAGCCAG
2101 CTAGAGGCCA TGATCCGCTT CCTCACCAGG AAGGACCCAG ACCTGGAGTG
2151 TGACGAGGAG CTGGTCCGGG AGGTGCTGTT TGACGCGGTG GTGACAGCCC
2201 CCATGGAAGC CTACTGGACA GCGCTGGCCC TCAACATGTC CGAGGAGTCT
2251 GAACACGTGG TGGACATGGC CTTCTGGGC ACCCGGGCTG GCCTCCTGAG
2301 AAGCAGCTTG TTCGTGGGCT CCGAGAAGGT CTCCGACAGG AAGTTCTCTGA
2351 CACCTGAGGA CGAGGCCAGC GTGTTCAACC TGGACCGCTT CCCGCTGTGG
2401 TACCGCCAGG CCTCAGAGCA TCCTGCTGGC AGCTTCGTCT TCAACCTCCG
2451 CTGGGCAGAA GGACCAGAAA GTGCGGGTGA ACCCATGGTG GTGACGGCAA
2501 GCACAGCTGT GCGGTGACC GTGGACAAGA GGACAGCCAT TGCTGCAGCC
2551 GCGGCGTCC AAATGAAGCT GGAATTCTC CAGCGCAAAT TCTGGGCGGC
2601 AACGCGGCAG TGCAGCACTG TGGATGGGCC GTACACACAG AGCTGCGAGG
2651 ACAGTGATCT GGACTGCTTC GTCATCGACA ACAACGGGTT CATTCTGATC
2701 TCCAAGAGGT CCCGAGAGAC GGAAGATTT CTGGGGGAGG TggaTGGTGC
2751 TGTCTTGACC CAGCTGCTCA GCATGGGGGT GTTCAGCCAA GTGACTATGT
2801 ATGACTATCA GGCCATGTGC AAACCTCGA GTCACCACCA CAGTGCAGCC
2851 CAGCCCCTGG TCAGCCCAAT TTCTGCCTTC TTGACGGCGA CCAGGTGGCT
2901 GCTGCAGGAG CTGGTGCTGT TCCTGCTGGA GTGGAGTGTC TGGGGCTCCT
2951 GGTACGACAG AGGGGCGgaG CCAAAAAGTG TCTTCCATCA CTCCCAAAA
3001 CACAagaagC AGGACCCGCT GCagcCTGc gacaCGgagt ACCccgTGt
3051 cGTGTAccaG CCGGccaTCC GGGaggCCAA CGGGATCGTG GAGTGCGGGC
```

3101 CCTGCCAGAA GGTATTTGTG GTGCAGCAGA TTCCCAACAG TAACCTCCTC  
 3151 CTCCTGGTGA CAGACCCAC CTGTGACTGC AGCATCTTCC CACCAGTGCT  
 3201 GCAGGAGGCG ACAGAAGTCA AATATAATGC CTCTGTCAAA TGTGACCGGA  
 3251 TGCGCTCCCA GaagctccGC CGGCGACCAG ACTCCTGCCA CGCCTTCCAT  
 3301 CCAGAGGTGC GGGTTGAGGC GGATCGAGGG TGGGCTGGAT TTTCATCCCC  
 3351 AAACCTCTG TGCCTGGGTC TGTGCCCCTG CAGACAGGAG CATATAGGGA  
 3401 TGCCAATGAA CACACCTGTG CCTGTGCTTC TCGGGGGAAA CATTGCGGTT  
 3451 TATGCCCTGT GACACTGTGA TATAATAAGA AACAGA

**Protein Sequence:**

M A V A L G T R R R D R V K L W A D T F G G D L Y N T  
 V T K Y S G S L L L Q K K Y K D V E S S L K I E E V D G L E  
 L V R K F S E D M E N M L R R K V E A V Q N L V E A A E E A  
 D L N H E F N E S L V F D Y Y N S V L I N E R D E K G N F V  
 E L G A E F L L E S N A H F S N L P V N T S I S S V Q L P T  
 N V Y N K D P D I L N G V Y M S E A L N A V F V E N F Q R D  
 P T L T W Q Y F G S A T G F F R I Y P G I K W T P D E N G V  
 I T F D C R N R G W Y I Q A A T S P K D I V I L V D V S G S  
 M K G L R M T I A K H T I T T I L D T L G E N D F V N I I A  
 Y N D Y V H Y I E P C F K G I L V Q A D R D N R E H F K L L  
 V E E L M V K G V G V V D Q A L R E A F Q I L K Q F Q E A K  
 Q G S L C N Q A I M L I S D G A V E D Y E P V F E K Y N W P  
 D C K V R V F T Y L I G R E V S F A D R M K W I A C N N K G  
 Y Y T Q I S T L A D T Q E N V M E Y L H V L S R P M V I N H  
 D H D I I W T E A Y M D S K L L S S Q A Q S L T L L T T V A  
 M P V F S K K N E T R S H G I L L G V V G S D V A L R E L M  
 K L A P R Y K L G V H G Y A F L N T N N G Y I L S H P D L R  
 P L Y R E G K K L K P K P N Y N S V D L S E V E W E D Q A E  
 S L R T A M I N R E T G T L S M D V K V P M D K G K R V L F  
 L T N D Y F F T D I S D T P F S L G A V L S R G H G E Y I L  
 L G N T S V E E G L H D L L H P D L A L A G D W I Y C I T D  
 I D P D H R K L S Q L E A M I R F L T R K D P D L E C D E E  
 L V R E V L F D A V V T A P M E A Y W T A L A L N M S E E S  
 E H V V D M A F L G T R A G L L R S S L F V G S E K V S D R  
 K F L T P E D E A S V F T L D R F P L W Y R Q A S E H P A G  
 S F V F N L R W A E G P E S A G E P M V V T A S T A V A V T  
 V D K R T A I A A A A G V Q M K L E F L Q R K F W A A T R Q  
 C S T V D G P Y T Q S C E D S D L D C F V I D N N G F I L I  
 S K R S R E T G R F L G E V D G A V L T Q Q L L S M G V F S Q  
 V T M Y D Y Q A M C K P S S H H S A A Q P L V S P I S A F  
 L T A T R W L L Q E L V L F L L E W S A V W G S W Y D R G A E  
 A K S V F H H S H K H K K Q D P L Q P C D T E Y P V F V Y Q  
 P A I R E A N G I V E C G P C Q K V F V V Q Q I P N S N L L  
 L L V T D P T C D C S I F P P V L Q E A T E V K Y N A S V K  
 C D R M R S Q K L R R R P D S C H A F H P E V R V E A D R G  
 W A G F S S P N P L C L G L C P C R Q E H I G M P M N T P V  
 P V L L G G N I R V Y A L

**Comparison of ha2δ-4.pep with ha2δ-3.pep:**

April 7, 2000 14:19

Percent Similarity: 69.172 Percent Identity: 60.133

1 .....MAVALGTRRRDR.....VKLWADTFG 21  
 : ||| | |||| ||  
 1 MAGPGSPRRASRGASALLAALLYAALGDVVRSEQQIPLSVVKLWASAFG 50  
 22 GDLYNTVTKYSGSLLLOKKYKDVESSLKIEEVDGLELVRKFSEDMENMLR 71  
 |: . ||||| |||||: | . |||: |||: ||: | . . . || |  
 51 GEIKSIAAKYSGSQLLOKKYKEYEKDVAIEEIDGLQLVKKLAKNMEEMFH 100

```

72 RKVEAVQNLVEAAEEADLNHEFNESLVFDYNSVLINERDEKGNFVELGA 121
:| |||. ||||| | |||. | :|:|. |||||. |||. |||
101 KKSEAVRRLVEAAEEAHLKHEFDADLQYEFNAVLINERDKDGNFLELGK 150

122 EFLLESNAHFSNLPVNTSISSVQLPTNVYNKDPDILNGVYMSEALNAV FV 171
||:| | ||. |||| | :| ||. |||. |||| | . |||| | |||. |||
151 EFILAPNDHFNNLPVNI SLSDVQVPTNMYNKDP AIVNGVYWSESLNKV FV 200

172 ENFQRDPTLTWQYFGSATGFFRIYPGIKWTPDENG VITFDCRNRGWY IQA 221
:| || |||. | ||||| |||| | ||||| ||||| ||||| |||||
201 DNFD RDPSLIWQYFGSAKGFFRQYPGIKWEPDENG VIAFDCRN RKWY IQA 250

222 ATSPKDIVILVDVSGSMKGLRMTIAKHTIT TILDTLGENDFVNIIAYNDY 271
|||||: ||||| ||||| : |||| | :.. ||||| :.. || |||||:
251 ATSPKDVILVDVSGSMKGLRLTIAKQTVSSILDTLGDDDFFNIIAYNEE 300

272 VHYIEPCFKGILVQADRDNREHF KLLVEELMVKG VGVVDQALREAFQ I LK 321
. ||: ||| | ||||| | : |||: . :. | ||: |.. | | ||| ||
301 LHYVEPCLNGTLVQADRTNKEHFREHLDKLF AKGIGMLDIALNEAFN ILS 350

322 QFQEAQQSGLCNQAIMLISDGAVEDYEPVFEKYNWPDCKVRVFTY LIGRE 371
| |||:|. |||||. ||||: | : :| ||||| |||: ||||| |||
351 DFNHTGQGSICSQAIMLITDGAVDTYDTIFAKYNWPD RKVRIFTY LIGRE 400

372 VSFADRMKWIACNNKGYTQISTLADTQENVMEYLHVLSRPMVINHDHDI 421
. ||| :|||. || |||: ||||| ||||| ||||| ||||| |||. :||:
401 AAFADNLKWMACANKGFFTQISTLADVQENVMEYLHVLSR PKVIDQEHDV 450

422 IWTEAYMSKLLSSQAQSLT..... LLTTVAMPVFSKKNETRSHGILLG 465
: |||||. || | ||| || | : ||||| |||||. |||| | |||||
451 VWTEAYIDSTL..PQAQKLTDQGPVLMTTVAMPVFSKQNETRSKGILLG 498

466 VVGSDVALRELMKLAPRYKLG VHG YAFLNTNNGYILSHPD LRPLYREGKK 515
|||. || .: ||: | | : ||||: ||||| ||||| |||. ||: || || |||
499 VVGTDVPVKELLKTIPKYKLG IGHYAFAITNNGYILTHPELRLLYBEGKK 548

516 LKPKPNYSVDLSEVEWEDQAESLRTAMINRETGTL SMDVKVPMDKGKRV 565
: |||||. ||||| |||||. : || ||: |||. || ||: || . |||||
549 .RRKPNYSVDLSEVEWEDRDDVLRNAMVNRTKGKFSMEVKKTVDKGKRV 597

566 LFLTNDYFFTDISDTPFSLGAVLSRGHGEYILLGNTSVEEGLHDL LHPDL 615
| : ||||: || ||||| ||||| |||||. || ||. : ||||| |||.
598 LVMTNDY YTDIKGTPFSLGVALSRGHGKYFFRGNVTIEEGLHDLEHPDV 647

616 ALAGDWIYCITDIDPDHRKLSQLEAMIRFLTRKDPDLECDEELVREVLFD 665
. || :| || ||: | :| |||||. :| | :| | :||. ||: . |||||
648 SLADEWSYCNTDLHPEHRHLSQLEAIKLYLKGKEPLLQCDKELIQEVLFD 697

666 AVVTAPMEAYWTALALNMSESEHVVDMAFLGTRAGLLRSSLFVGSEKVS 715
|||. ||. |||||. |||| | | : | :. ||||| || | . |||||. |...
698 AVVSAPIEAYWTSALANKSENSDKGVEVAFLGTRTGLSRINL FVGAEQLT 747

716 DRKFLTPEDEASVFTLDRFPLWYRQASEHPAGSFVFNLRWAEGPESAGEP 765
.. || |. :| | |||||. |. | ||||:.. .. ||
748 NQDFLKAGDKENIFNADHFPLWYRRAAEQIPGSFVYSIPFSTGP..VNKS 795

766 MVVTASTAVAVTVDKRTAIAAAAAGVQMKLEFLQRKFWAATRQCSTVDGPY 815
|||||: . . :.. : || | : ||||| ||||| |. |||. |... ||
796 NVVTASTSIQLLDERKSPVVAAGVIQMKLEFFQRKFWTASRQCASLDGKC 845

816 TQSCEDSDLD CFVIDNNGFILISKRSRETGRFLGEVDGAVLTQLLSMGVF 865
. ||:| ..:|. ||||| |||:|. :|| | ||: |||: . ||. || |

```

846 SISCDDDETVCYLIDNNGFILVSEDYQTGTGDFGEIEGAVMNKLLTMGSF 895  
 866 SQVTMYDYQAMCKPSSHHSAAQPLVSPISAFILTATRWLLQELVLFLEW 915  
 .:|:|||||: . | . | .|||.| :|:| |||||.|. |.  
 896 KRITLYDYQAMCRANKESSDGAHGLLDPYNAFLSAVKWIMTELVLFLVEF 945  
 916 SVWGSWYDRGAEAKSVFHSHKHKKQDPLQPCDTEYPVFVYQPAIREANG 965  
 .. ||: ||. | . |:||||||| || : |:| |  
 946 NLC.SWWHSDMTAKA.....QKLKQTLPCDTEYPAFVSERTIKETTG 987  
 966 IVECGPCQKVVFVQQIPNSNLLLLVTDPTCDCSIFPPVLQEATEVKYNAS 1015  
 : | | | |:||||.||| :. | | . | | : | :| |  
 988 NIACEDCSKSFVIQQIPSSNLFMVVVDSSCLCESVAPITMAPIEIRYNES 1037  
 1016 VKCDRMRSQKLRRRPDSCHAFHPEVRVEADRGWAGFSSPN.....PLCLG 1060  
 .||:|:..||:||||:|||| | . || |  
 1038 LK CERLKAQKIRRRPESCHGFHPEENARECGGAPSLQAQTVLLLLPLLLM 1087  
 1061 LCPCRQEHIGMPMNTFVPVLLGGNIRVYAL 1090  
 |  
 1088 LFSR..... 1091

**This Page is Inserted by IFW Indexing and Scanning  
Operations and is not part of the Official Record**

**BEST AVAILABLE IMAGES**

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images include but are not limited to the items checked:

- ☐ **BLACK BORDERS**
- ☐ **IMAGE CUT OFF AT TOP, BOTTOM OR SIDES**
- ☐ **FADED TEXT OR DRAWING**
- ☐ **BLURRED OR ILLEGIBLE TEXT OR DRAWING**
- ☐ **SKEWED/SLANTED IMAGES**
- ☐ **COLOR OR BLACK AND WHITE PHOTOGRAPHS**
- ☐ **GRAY SCALE DOCUMENTS**
- ☐ **LINES OR MARKS ON ORIGINAL DOCUMENT**
- ☐ **REFERENCE(S) OR EXHIBIT(S) SUBMITTED ARE POOR QUALITY**
- ☐ **OTHER:** \_\_\_\_\_

**IMAGES ARE BEST AVAILABLE COPY.**

**As rescanning these documents will not correct the image problems checked, please do not report these problems to the IFW Image Problem Mailbox.**